

CLAIMS AS AMENDED UNDER ART.19 PCT

- 1) Circular recombinant plasmid DNA construct encoding a) a protein tag, b) a visual marker protein, and containing c) a multiple cloning site suitable for insertion of an additional gene, characterised in that the gene sequence encoding the protein tag and the visual marker protein are specifically designed and engineered at the DNA level for respectively a) immobilisation purposes and b) visualisation and quantification purposes at the protein level.
- 2) Construct according to claim 1, characterised in that the protein tag is chosen from the group containing lysine (lys), histidine (his), tyrosine (tyr), phenylalanine (phe), arginine (arg), glutamic acid (glu), aspartic acid (asp), glutamate, aspartate, asparagine (asn), glycine (gly), glutamine (gln), alanine (ala), valine (val), tryptophan (trp).
- 3) Construct according to claim 1 or 2, characterised in that the protein tag is a histidine-tag such as a polyhistidine variant, in particular (6X) histidine.
- 4) Construct according to any one of the preceding claims, characterised in that the visual marker protein is chosen from the group containing fluorescent or phosphorescent proteins, wherein the fluorescent protein is chosen from the group containing Green Fluorescent Protein (GFP), Red Fluorescent Protein (RFP), Yellow Fluorescent Protein (YFP) and Blue Fluorescent Protein (BFP) as well as their variants and/or mutants.
- 5) Construct according to any one of the preceding claims, characterised in that the multiple cloning site contains restriction enzyme recognition sites chosen from the group containing SacI, Sal I, Hid III, Eag I, Not I.
- 6) Construct according to any one of the preceding claims, characterised in that it further contains a frame adapter of variable length between the visual marker and protein tag genes.
- 7) Construct according to any one of the preceding claims, characterised in that it expresses a fusion protein, wherein the tag is suitable to interact directly with appropriate surface pendant groups of a support material.
- 8) Construct according to claim 9, characterised in that the direct interaction with the support material is covalent or non-covalent.
- 9) Construct according to claim 10, characterised in that the direct interaction is non-covalent and yet freely accessible and leach-free like covalent one.

- 10) Method for preparing and immobilising a protein on a support material, characterised in that it contains the steps of:
- a) Engineering at the DNA level, in series a protein tag suitable to interact directly with appropriate surface pendant groups of a support material, a fluorescent marker protein for visualisation and quantification purposes at the protein level and a multiple cloning site suitable for insertion of a target protein to be immobilised,
 - b) Inserting the corresponding gene of the target protein to be immobilised into the multiple cloning site;
 - c) Initiating protein expression.
 - d) Optionally pre-treating the support material;
 - e) Incubating the protein and support material together, wherein the protein is immobilised to the support via specific tag-surface interactions;
 - f) Washing away the non-specific biomolecules;
 - g) Optionally quantifying the fluorescence of the visual marker protein;
 - h) Optionally desorbing the target protein.
- 11) Method according to claim 10, characterised in that the support material is chosen from the group containing polymers, biopolymers, glass and composites containing silicone dioxides, metals and metal oxides, as well as any combination thereof on the microscopic, mesoscopic or macroscopic length scale.
- 12) Method according to any one of claims 10 and 11, characterised in that the support material is chosen from the group containing polymers, silicon dioxides, aluminum oxides, titanium oxides, magnesium oxides, borates, metals and other metal oxides.
- 13) Method according to any one of claims 11 and 12, characterised in that the polymers are chosen from the group containing polyolefins such as polystyrene, polyacrylates, polymethacrylates, polybutylene, polyvinylalcohol and related derivatives, polyvinylchlorides, polyisoprene, polypropylene, polyphenols, polyamides, polyesters polysulfones, polyethersulfones, polyethersulfides, polyimines, polyethyleneglycols, polypropyleneglycols, polyimides, polycarbonates, polyurethanes,

- 5 14) Method according to claim 13, characterised in that the polymer surface is chemically treated to bear various functional groups chosen between carboxyl groups, hydroxyl groups, amino groups, amide groups, ester groups, imide groups, imine groups, mercapto groups, nitro groups, sulfonate groups, phosphate groups, phosphonate groups, cyano groups, sulfone groups, aldehyde groups, epoxide groups, urethane groups, ketone groups, phenolic groups, aromatic groups, alkyl, alkenyl, alkynyl, acyl and aryl groups, silanol groups, silicon oxide groups, siloxane groups, metal hydroxide groups, metal oxide groups, and elemental metals.
- 10 15) Method according to any one of claims 13 to 16, characterised in that the support material is carboxylated polystyrene.
- 15 16) Method according to claim 10, characterised in that quantifying in step g) of the fluorescence of the visual marker protein is used in applications selected from analysis, diagnosis (like in enzyme based diagnostic kits), incubation, storage, sensing, arraying and orienting, catalysis, stabilisation, binding, signal transduction, chemical transformation, implant passivation and surface biocompatibilization, surface activation, purification, detoxification and scavenging.

**[Received by the International Bureau on 04 DEC 2003 (04.12.03) ;
original claims 1 to 3, unchanged ; original claims 4 to 16, amended ;
claims 17 to 20, cancelled]**

- 1) Circular recombinant plasmid DNA construct encoding a) a protein tag,
b) a visual marker protein, and containing c) a multiple cloning site suitable for
insertion of an additional gene, characterised in that the gene sequence encoding
the protein tag and the visual marker protein are specifically designed and
engineered at the DNA level for respectively a) immobilisation purposes and
b) visualisation and quantification purposes at the protein level.
- 2) Construct according to claim 1, characterised in that the protein tag is chosen from
the group containing lysine (lys), histidine (his), tyrosine (tyr), phenylalanine (phe),
arginine (arg), glutamic acid (glu), aspartic acid (asp), glutamate, aspartate,
asparagine (asn), glycine (gly), glutamine (gln), alanine (ala), valine (val),
tryptophan (trp).
- 3) Construct according to claim 1 or 2, characterised in that the protein tag is a
histidine-tag such as a polyhistidine variant, in particular (6X) histidine.
- 4) Construct according to any one of the preceding claims, characterised in that the
visual marker protein is chosen from the group containing fluorescent or
phosphorescent proteins, wherein the fluorescent protein is chosen from the group
containing Green Fluorescent Protein (GFP), Red Fluorescent Protein (RFP),
Yellow Fluorescent Protein (YFP) and Blue Fluorescent Protein (BFP) as well as
their variants and/or mutants.
- 5) Construct according to any one of the preceding claims, characterised in that the
multiple cloning site contains restriction enzyme recognition sites chosen from the
group containing SacI, Sal I, Hid III, Eag I, Not I.
- 6) Construct according to any one of the preceding claims, characterised in that it
further contains a frame adapter of variable length between the visual marker and
protein tag genes.
- 7) Construct according to any one of the preceding claims, characterised in that it
expresses a fusion protein, wherein the tag is suitable to interact directly with
appropriate surface pendant groups of a support material.
- 8) Construct according to claim 9, characterised in that the direct interaction with the
support material is covalent or non-covalent.
- 9) Construct according to claim 10, characterised in that the direct interaction is non-
covalent and yet freely accessible and leach-free like covalent one.

10) Method for preparing and immobilising a protein on a support material, characterised in that it contains the steps of:

- a) Engineering at the DNA level, in series a protein tag suitable to interact directly with appropriate surface pendant groups of a support material, a fluorescent marker protein for visualisation and quantification purposes at the protein level and a multiple cloning site suitable for insertion of a target protein to be immobilised,
- b) Inserting the corresponding gene of the target protein to be immobilised into the multiple cloning site;
- c) Initiating protein expression.
- d) Optionally pre-treating the support material;
- e) Incubating the protein and support material together, wherein the protein is immobilised to the support via specific tag-surface interactions;
- f) Washing away the non-specific biomolecules;
- g) Optionally quantifying the fluorescence of the visual marker protein;
- h) Optionally desorbing the target protein.

11) Method according to claim 10, characterised in that the support material is chosen from the group containing polymers, biopolymers, glass and composites containing silicone dioxides, metals and metal oxides, as well as any combination thereof on the microscopic, mesoscopic or macroscopic length scale.

12) Method according to any one of claims 10 and 11, characterised in that the support material is chosen from the group containing polymers, silicon dioxides, aluminum oxides, titanium oxides, magnesium oxides, borates, metals and other metal oxides.

13) Method according to any one of claims 11 and 12, characterised in that the polymers are chosen from the group containing polyolefins such as polystyrene, polyacrylates, polymethacrylates, polybutylene, polyvinylalcohol and related derivatives, polyvinylchlorides, polyisoprene, polypropylene, polyphenols, polyamides, polyesters polysulfones, polyethersulfones, polyethersulfides, polyimines, polyethyleneglycols, polypropyleneglycols, polyimides, polycarbonates, polyurethanes,

14) Method according to claim 13, characterised in that the polymer surface is chemically treated to bear various functional groups chosen between carboxyl groups, hydroxyl groups, amino groups, amide groups, ester groups, imide groups, imine groups, mercapto groups, nitro groups, sulfonate groups, phosphate groups, phosphonate groups, cyano groups, sulfone groups, aldehyde groups, epoxide groups, urethane groups, ketone groups, phenolic groups, aromatic groups, alkyl, alkenyl, alkynyl, acyl and aryl groups, silanol groups, silicon oxide groups, siloxane groups, metal hydroxide groups, metal oxide groups, and elemental metals.

15) Method according to any one of claims 13 to 16, characterised in that the support material is carboxylated polystyrene.

16) Method according to claim 10, characterised in that quantifying in step g) of the fluorescence of the visual marker protein is used in applications selected from analysis, diagnosis (like in enzyme based diagnostic kits), incubation, storage, sensing, arraying and orienting, catalysis, stabilisation, binding, signal transduction, chemical transformation, implant passivation and surface biocompatibilization, surface activation, purification, detoxification and scavenging.

IN THE CLAIMS

1. (Previously Presented) Circular recombinant plasmid DNA construct encoding a) a protein tag, b) a visual marker protein, and containing c) a multiple cloning site suitable for insertion of an additional gene, characterised in that it further contains d) a frame adaptor of variable length between the visual marker and protein tag genes and in that the gene sequence encoding the protein tag, the visual marker protein and the frame adaptor are specifically designed and engineered at the DNA level for respectively i) immobilisation purposes, ii) visualisation and quantification purposes at the protein level, and iii) providing a large distance separating protein and surface to enable the immobilised enzymes to display native-like characteristics.

2. (Previously Presented) Construct according to claim 1, characterised in that the protein tag is chosen from the group containing lysine (lys), histidine (his), tyrosine (tyr), phenylalanine (phe), arginine (arg), glutamic acid (glu), aspartic acid (asp), glutamate, aspartate, asparagine (asn), glycine (gly), glutamine (gln), alanine (ala), valine (val), tryptophan (trp).

3. (Currently Amended) Construct according to claim 1 or 2, characterised in that the protein tag is a histidine-tag such as a polyhistidine variant, in particular (6X) histidine.

4. (Currently Amended) Construct according to ~~any one of the preceding claims~~ claim 1, characterised in that the visual marker protein is chosen from the group containing fluorescent or phosphorescent proteins, wherein the fluorescent protein is chosen from the group containing Green Fluorescent Protein (GFP), Red Fluorescent Protein (RFP), Yellow Fluorescent Protein (YFP) and Blue Fluorescent Protein (BFP) as well as their variants and/or

mutants.

5. (Currently Amended) Construct according to ~~any one of the preceding claims~~ claim 1, characterised in that the multiple cloning site contains restriction enzyme recognition sites chosen from the group containing SacI, Sal I, Hind III, Eag I, Not I.

6. (Currently Amended) Construct according to ~~any one of the preceding claims~~ claim 1, characterised in that it expresses a fusion protein, wherein the tag is suitable to interact directly with appropriate surface pendant groups of a support material.

7. (Previously Presented) Construct according to claim 6, characterised in that the direct interaction with the support material is covalent or non-covalent.

8. (Previously Presented) Construct according to claim 7, characterised in that the direct interaction is non-covalent and yet freely accessible and leach-free like covalent one.

9. (Currently Amended) Method for preparing and immobilising a protein on a support material, comprising:

a) Engineering at the DNA level, in series a protein tag suitable to interact directly with appropriate surface pendant groups of a support material, a fluorescent marker protein for visualisation and quantification purposes at the protein level and a multiple cloning site suitable for insertion of a target protein to be immobilised,

b) Inserting the corresponding gene of the target protein to be immobilised into the multiple cloning site;

c) Initiating protein expression.

- d) Optionally pre-treating the support material;
- e) Incubating the protein and support material together,
wherein the protein is immobilised to the support via specific tag-surface interactions;
- f) Washing away the non-specific biomolecules;
- g) Optionally quantifying the fluorescence of the visual
marker protein;
- h) Optionally desorbing the target protein[. .].

characterised in that the support material is chosen from the group containing polymers, biopolymers, glass and composites containing silicone dioxides, metals and metal oxides, as well as any combination thereof on the microscopic, mesoscopic or macroscopic length scale.

10. (Previously Presented) Method according to claim 9, characterised in that the support material is chosen from the group containing polymers, silicon dioxides, aluminum oxides, titanium oxides, magnesium oxides, borates, metals and other metal oxides.

11. (Currently Amended) Method according to ~~any one of claims~~ claim 9 and 11, characterised in that the polymers are chosen from the group containing polyolefins such as polystyrene, polyacrylates, polymethacrylates, polybutylene, polyvinylalcohol and related derivatives, polyvinylchlorides, polyisoprene, polypropylene, polyphenols, polyamides, polyesters polysulfones, polyethersulfones, polyethersulfides, polyimines, polyethyleneglycols, polypropyleneglycols, polyimides, polycarbonates, polyurethanes.

12. (Previously Presented) Method according to claim 11, characterised in that the polymer surface is chemically treated to bear various functional groups chosen between carboxyl groups, hydroxyl groups, amino groups, amide groups, ester groups, imide groups, imine groups, mercapto groups, nitro groups, sulfonate groups, phosphate groups, phosphonate

groups, cyano groups, sulfone groups, aldehyde groups, epoxide groups, urethane groups, ketone groups, phenolic groups, aromatic groups, alkyl, alkenyl, alkynyl, acyl and aryl groups, silanol groups, silicon oxide groups, siloxane groups, metal hydroxide groups, metal oxide groups, and elemental metals.

13. (Currently Amended) Method according to ~~any one of claims~~ claim 9 to 12, characterised in that the support material is carboxylated polystyrene.

14. (Previously Presented) Method according to claim 9, characterised in that quantifying in step g) of the fluorescence of the visual marker protein is used in applications selected from analysis, diagnosis (like in enzyme based diagnostic kits), incubation, storage, sensing, arraying and orienting, catalysis, stabilisation, binding, signal transduction, chemical transformation, implant passivation and surface biocompatibilization, surface activation, purification, detoxification and scavenging.

15. (Currently Amended) A two-component system obtained by ~~any one of claims~~ claim 9 to 14.

16. (Currently Amended) A two-component system according to claim 15, described by and an activated support material and a protein encoded by a recombinant plasmid DNA construct ~~according to any one of claims 1 to 8~~ encoding a) a protein tag, b) a visual marker protein, and containing c) a multiple cloning site suitable for insertion of an additional gene, characterised in that it further contains d) a frame adapter of variable length between the visual marker and protein tag genes and in that the gene sequence encoding the protein tag, the visual marker protein and the frame adaptor are specifically designed and engineered at

the DNA level for respectively i) immobilisation purposes, ii) visualisation and quantification purposes at the protein level, and iii) providing a large distance separating protein and surface to enable the immobilised enzymes to display native-like characteristics.